

**Amendments to the Specification:**

Please replace the last paragraph on page 6 with the following amended paragraph:

A representation of the nucleotide sequence of Clone 20468752.0.18 is shown in Table 2 and includes a nucleotide sequence (SEQ ID NO:1) of 1867 bp. This nucleotide sequence has an open reading frame (ORF) encoding a polypeptide of 567 amino acid residues (SEQ ID NO:2) with a predicted molecular weight of 63327 Daltons. The start codon is located at nucleotides 128-130 and the stop codon is located at nucleotides 1829-1831. The protein encoded by Clone 20468752.0.18 (SEQ ID NO:2) was predicted by the PSORT program to be extracellularly localized with a certainty of 0.3700. Analysis using the PSORT and SignalP computer programs predicted that there is ~~may be~~ a signal peptide with the most likely cleavage occurring between residues 21 and 22, at the sequence ISS-LP. The nucleic acid (SEQ ID NO:1) and amino acid (SEQ ID NO:2) sequences of Clone 20468752.0.18 is shown below in Table 2.

Please replace the last paragraph on page 9 with the following amended paragraph:

The protein (SEQ ID NO:4) encoded by Clone 20468752.0.18-U is predicted by the PSORT program to extracellularly localized with a certainty of 0.3700. Analysis with the PSORT and SignalP computer programs predicted that there is ~~may be~~ a signal peptide, with the most likely cleavage occurring between residues 21 and 22, at the sequence ISS-LP. The nucleic acid (SEQ ID NO:3) and amino acid (SEQ ID NO:4) sequences of Clone 20468752.0.18-U is shown below in Table 3.